OIPE

Does Not Campy Correcte 11 Shorte No. 1

RAW SEQUENCE LISTING

PATENT APPLICATION US/09/676,475

PATE: 10/11/12/20 TIME: 18:21,53

Input Set : A:\LJ4377.app
Output Set : N:\CRF3\10122000\1676475.raw

3 -116 - APPLICANT: Bugslahti, Erkki Pasqualini, Renara

Eaglite Daniel

12-- Time OF invention: Nethods of Identifying Lang Homing Molecules Using Mombrane Dipeptidate 13 - 137 - FILE AVFERENCE: P-1A 12 <140 > CURRENT APPLICATION NUMBER: US/09/676,475 13 <141 > CURRENT FILING DATE: 2000-09-29 13 | 150 - REFOR APPLICATION NUMBER: 09/542.467 15 -151 - PRIOR FILING DATE: 1968 03:13 13 - 160 - NUMBER OF SEQ ID NOS: 152 20 .170 - SOFTWARE: ParentIn Ver. 2.0 22 -210 - STO ID NO: 1 23 -211 - LENGTH: 13 TOURS PRAIDRE: 3 - 229 - CHER INFORMATION, Dobbi prion of Artificial Sequence: Symbatic 0 - 150 - 21 20 NOE: 1 .1 Cys GI: Phe Glu Cys Val Arg Gln Cys Pro Glu Arg Cys 12 1 15 - 210 - SEQ ID NO: 2 15 -211 - LENGTH: 8 222 - INPE: PPT 213 - OPCANISM: Artificial Sequence 10 120 FEA.URE: 10 200 FEA.URE: 10 200 FEA.URE: 10 200 FEA.URE: 11 200 FEA.URE: 12 200 SECONOCE: 2 11 200 Gly Pho Glu Leg Gir Tin Spa 48 210 - SEQ ID NOT 3 49 211: LENGTH: 13 50:-212: TYPE: PRT 51:-213: OFGANISM: Artificial Sequence \$3 - 200 - FRATURE: 14 - 1223 - OTHER INFORMATION, Description of Artificial Sequence, Synthetic \$6 - 103 - SEQUENCE: 3 81 | 119 | SPQ LD NO: 1 82 | 119 | LENGTH: 9 63 | 210 | TYPE: PRI 64 | 213 | ORGANISM: Artificial Sequence 86 - 220 - FEATURE: 67 - 223 - OTHER INFORMATION: Description of Artificial Sequence: Synthetic 69 - 100 - SEQUENCE: 1 TO Set Tip dys Gld Pro Gly Tip dys Arg

Q.5

RAW SEQUENCE LISTING DATE: 10/12/2000 FAIENT APPLICATION: US/09/676,475 Fifth: 18.27:53

Input Set : A:\LJ4377.app
tutput Set: N:\CRF3\10122000\1676475.raw

```
71 - 210: SEQ ID NO: 3
^{15} \pm 2115 | LENGTH \pm 9
T6 - 1125 TYPE: PPT
     .213 * ORGANISM: Artificial Jequence
79 220% FEATURE:
80 223 OTHER INFORMATION: Description of Artificial Sequence: Synthetic
62 - 100% SEQUENCE: 5
83 Cys Ser Cys Phe Ard Asp Val Cys Cys
87 <2105 SEQ ID NO: 6
88 <2115 LENGIH: 9
89 SE125 TYPE: PRT
%0 -213 - ORDANISM: Artificial Sequence
90 - 320 - FEATURE:
93 #323> OTHER INFORMATION: Description of Artitional Sequence: Southetic
95 - 400> SEQUENCE: 6
36 Cys Aig Asp Val Val Ser Val Ile Cys
9° 1 5

106 -219 -880 IT NO. "

101 -210 LENSTH: "

103 -2220 TUPE PRT

103 -2230 -06GPNISM: Artificial Sequence

105 -2230 THEP INFOSMATION: Description of Artificial Sequence: Spatiet!"

106 -2230 THEP INFOSMATION: Description of Artificial Sequence: Spatiet!"

106 -4100 SEQUENCE: "

107 Jan San Pro Pro Tor Met Tin
. H Leu Ser Pro Pro Tyr Met Tip
1..0
1.3 <210 > SEQ ID NO: 8
La solle Length: 8
1.5 \times 212\% Type: PRT 1.8 \times 213\% GRGANISM: Artificial Sequence
128 -220> FEATURE:
i 6 s223: STHER INFORMATION: Description of Artificial Sequence. Symbotic 1.21 \pm 400 \times SEQUENCE: 8
122 Gly fle Gly Glu Val Glu Val Cys
123
176 210 SEQ IP NO: 9
127 (211> LENGTH: 7
128 -212 - TYPE: PRI
129 - 2132 OPGANISM: Artificial Sequence
141 42205 FEATURE:
182 8223 OTHER INFORMATION: Description of Artificial Sequence. Synthetic
134 <400 - SEQUENCE: 9
195 Tyr Ser Gly Lys Trp Gly Lys
1.6
139 <210 - SEQ ID NO: 10
140 -211: LENGTH:
141 <212> TYPE: PFT
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 RAW SEQUENCE LISTING
 DATE: 10/1:/2000

 PATENT APPLICATION:
 US/09/676,475
 TIME: 18:27:53

Input Set : A:\LJ4377.app
Output Set: N:\CRF3\10122000\1676475.raw

```
140 <213> ORGANISM: Artificial Sequence
144 <2205 FEATURE
115 <2232 OTHER INFORMATION: Description of Artificial Sequence: Synthetic 117 <4008 SEQUENCE: 10 116 GJu Val Arg Ser Arg Lou Ser
1.454
152 32108 SE2 ID NO: 11
152 32108 SE2 ID NO: 11
153 32128 LENGTH: 7
151 32128 TYPE: PRT
15% 213% ORGANISM: Artificial Sequence
15% 220% FEATURE
158 3235 OTHER INFORMATION: Description of Artificial Sequence: Synthetic
146 \sim 100 > \text{SEQUENCE}: 11
183, Arg Val 31y teu Val Ala Arg
1.6 %1.00 SEQ 1D NO: 12
166 %1.10 LENGTH: 7
167 %1.00 TYPE: PRT
198 tl. % ORGANISM. Artificial Sequence
1 % CLOS FRATURE
11. 32000 OTHER INFORMATION: Description of Artificial Sequence: Synthetic
    c.00. SEQUENCE: 12
1 4 Val Lys Ser Val Cys Arg Thr
1 & <11(> SEQ ID NO: 13
1 9 <211 : LENGTH: 9
I-( -2115> TYPE: PRT
141 - 21 -> GRGANISM: Artificial Sequence
1 1 KIPOS FEATURE
1.4 <10 s OTHER INFORMATION: Description of Artificial Sequence: Synthetic 1.6 <400 SEQUENCE: 13
1.7 Cys Let Ala Lys Giu Asn Val Val Cys
1...8
191 Paul09 SEQ 15 NO: 14
191 d2.10 LENGTH: 8
195 d212> TYPE: PRT
194 ×215> CRGANISM: Artificial Sequence
196 ×216> FEATURE:
155 -1139 CTHEF INFORMATION: Description of Artificial Sequence: Synthetic 155 -4409 SEQUENCE: 14\,
2) ( Cys Gly Gle Phe Lys Val Gly Cys
3 | 1
201 0110> SEQ ID NO: 15
20% 3211> LENGIH: 8
208 3712> IYPE: PRT
207 <213> ORGANISM: Artificial Sequence
209 - 220> FEATURE:
210 <221> CTHER INFOFMATION: Description of Artificial Sequence: Synthetic
212 <400 > SEQUENCE: 15
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DATE: 1071172000
TIME: 18:27:63
                        PATENT APPLICATION: US/09/676,475
                        Input Set : A:\LJ4377.app
                        Output Set: N:\CRF3\10122000\1676475.raw
    213 Cys Thr Leu Arg Asp Arg Asn Cys
    211
         1 3
+ 216 + SEQ ID NO: lo
   217 (216 - SEQ ID No: 10
213 - 211 - IENGTH: 8
214 - 212 - TYPE: PRI
224 - 213 - GROANISM: Artificial Sequence
224: 8229 - FEATURE:
227 - 2230 OTHER INFORMATION: Description
    203 - 2235 OTHER INFORMATION: Description of Artificial Sequence: Synthetic ads + 1005 SEQUENCE: 16
    \pm 2\sigma Cys Ile Gly Glu Vai Glu Vai Cys
   130 100 SEQ 10 No: 17
25. dir Length: 28
152 212 TYPE, PRT
153 212 GREANISM: Artificial Sequence
155 226 FEATURE:
    Plo Bully FENISHE:
   .1. 2019 NAME/KET: SITE
[13] -2x1 - Location: (11)...(23)
[11] -2r12 Other Information: Kaa may be present or absent.
   .19 .230 FEATURE: Car - 223 - OTHER INFORMATION: Description of Artificial Sequence: Synthetic
   218 - 100 - SEQUENCE: 1'
250 1
>> 252 Xaa Xaa Xaa Xaa Xaa Xaa Xaa
                        7.0
   DIS 20 DIS SEQ IF NO: 18

FOR This LEMOTH: T

FOR DIS TYPE: IPI

FOR DIS ORGANISM: Artificial Sequence
   26. 220. FEATURE: 26.2 - 223. OTHER INFORMATION: Description of Artificial Sequence: Synthetic
    26: - 190 > SEQUENCE: 18
   Poly Lou Cys Thr Ala Met Thr Glu
   JAL.
   200 - 210 - SEQ ID NOTE LO
   In - 111 = LENGTH: 8
   27.1 > 21.2 > TASE: DEL
         213 ORGANISM: Artificial Sequence
    271 × 220 FEATUPE:
    27 - 223 - OTHER INFORMATION: Description of Artificial Sequence: Synthetic
         1000 SEQUENCE: 19
    1.16 GPy Ile Ser Ala Leu Val Leu Ser
   283 310 × SEQ ID No: 20
```

RAW SEQUENCE LISTING

283 +211 - LENGTH: 8 281 -213 - TYPE: PRI 28% $\sim 213 \times 0.09$ GANISM: Artificial Sequence 28% $\sim 220 \times \text{FFATURE}$: THE CLUST GIRP INFORMATION: Description of Artificial Sequence: Synthetic for $+100 - \rm cm/\, \rm TeV$. 79. Cys Gly 1/s Arg 1/1 Arg Ash C/s 29 + 5210, SEQ 1D NO: 21 096 -211 FindTR. 097 -212 TYPE: PRT 098 -213 / PGAMISM: Artificial Sequence 100 - 210 - FEATUPE 301-223 . These insummation: London prior of Arbificial Sequence: Synthetic 30×1400 - SPQUENCE- 21:04 Ser Mot Ser Ile Ala Arg Leu c2i0 - FEATURE: 14 - 22% WIHER INFORMATION: Description of Artificial Sequence: Synthetic te sime seguence . i Val Sor The Lei Gla Tyr Arg 11 -210 - SFQ 1E NO: 23 -22 -211 - LENGIH: 9 -22 S212 - TYPE: PRI -24 S213 - OPGANISM: Artificial Sequence 16 1980 THATURE: 10 - 831 NAMEZKEY: UNSUPK $(21 \pm 222 \pm 13) \text{CATION} \colon (21) \colon (2)$ +12 (27), GIRER INFORMATION: Unmile 11 (210) FRATURE: 32 (221) NAME/FEU UNIVER: (30) (222) INVATION: (1) (1) -) Cys is at location 8. Do you mean 9? -2004 OTHER INFORMATION: Unsure 39 FORD FRATURE: 10 .201 NAMEYEY: UNSUPE 11 .02 LOCATION (8) 12 .113 (THER INFERENCE) This Pro 14 .1000 SEQUENCE: 13 W--> 345 Xaa Xaa Cys Xaa Xaa Xaa Xaa Cys (Xaa . 16 1

DAFE: 10/12/2000

TIME: 18:27:53

RAW SEQUENCE LISTING

PATENT APPLICATION: US, 09/676, 475

FYI

349 (210 - 802 to NO: 34 350 (21) - LENGIE: 2

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

VERIFICATION SUMMARY

RATENI APPLICATION US/09/676,475

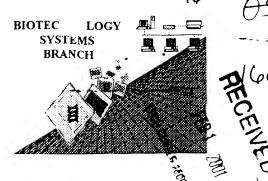
DATE: 20/12/2000 IIME: 18:27.:4

Input Set A:\LJ4377.app

Output Set: N:\CRF3\10122000\1676475.raw

Lili Mich C: Current Application Number differs, Peplaced Application Number Lili Mich C: Current Filing Late differs, Replaced Carrent Filing Date Lilis Mich, W: (10, "a" an "saa" used, for SEQ 10=.1 [1-252 Mich] W: (10) "r" or "saa" used, for SEQ 10=.1 [1-252 Mich] W: (10) "n" or "saa" used, for SEQ 10=.1 [1-252 Mich] W: (10) "n" or "saa" used, for SEQ 10=.2 [1-252 Mich] W: (10) "n" or "saa" used, for SEQ 10=.2 [1-252 Mich] W: (10) "n" or "saa" used, for SEQ 10=.2 [1-252 Mich] W: (10) "n" or "saa" used, for SEQ 10=.2 [1-252 Mich] W: (10) "n" or "saa" used, for SEQ 10=.2 [1-252 Mich] W: (10) "n" or "saa" used, for SEQ 10=.2 [1-252 Mich] W: (10) "n" or "saa" used, for SEQ 10=.3 [1-252 Mi

RAW SEQUENCE LISTING ERROR REPORT



The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 09/676,475

OIPE

Date Processed by STIC:

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS. PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,

TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216. PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax) PATENTIN 3.0 e-mail help: patin30help@uspto.gov or phone 703-306-4119 (R. Wax)

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 3.0 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW:

Checker Version 3.0

The Checker Version 3.0 application is a state-of the-art Windows based software program employing a logical and intuitive user-interface to check whether a sequence listing is in compliance with format and content rules. Checker Version 3 0 works for sequence listings generated for the original version of 37 CFR §§1.821 – 1.825 effective October 1, 1990 (old rules) and the revised version (new rules) effective July 1, 1998 as well as World Intellectual Property Organization (WIPO) Standard ST.25

Checker Version 3.0 replaces the previous DOS-based version of Checker, and is Y2Kcompliant. Checker allows public users to check sequence listings in Computer Readable form (CRF) before submitting them to the United States Patent and Trademark Office (USPTO). Use of Checker prior to filing the sequence listing is expected to result in fewer errored sequence listings, thus saving time and money.

Checker Version 3.0 can be down loaded from the USPTO website at the following address: http://www.uspto.gov/web/offices/pac/checker